# **Resource Summary Report**

Generated by dkNET on Apr 16, 2025

# **PRESTO: Genetic Association Analysis Software**

RRID:SCR\_013285

Type: Tool

### **Proper Citation**

PRESTO: Genetic Association Analysis Software (RRID:SCR\_013285)

#### Resource Information

**URL:** http://faculty.washington.edu/browning/presto/presto.html

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**Description:** Software application that performs permutation testing and computes empirical distributions of order statistics for one and two stage association studies with stratified or unstratified data.

**Resource Type:** software resource, software application

**Defining Citation:** DOI:10.1093/bioinformatics/btu138

**Keywords:** gene, genetic, genomic, java, ms-windows, unix, solaris, linux, macos

**Funding:** 

Resource Name: PRESTO: Genetic Association Analysis Software

Resource ID: SCR\_013285

Alternate IDs: nlx\_154549

**Record Creation Time:** 20220129T080315+0000

**Record Last Update:** 20250416T063639+0000

## Ratings and Alerts

No rating or validation information has been found for PRESTO: Genetic Association Analysis Software.

No alerts have been found for PRESTO: Genetic Association Analysis Software.

#### Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at <u>dkNET</u>.

Rubelt F, et al. (2016) Individual heritable differences result in unique cell lymphocyte receptor repertoires of naïve and antigen-experienced cells. Nature communications, 7, 11112.